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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/985,675

DATE: 02/22/2002

TIME: 09:00:06

Ala Gly

Input Set : N:\Crf3\RULE60\09985675.raw Output Set: N:\CRF3\02222002\I985675.raw

SEQUENCE LISTING

```
(1) GENERAL INFORMATION:
               (i) APPLICANT: CROSIER, PHILIP S.
        6
        7
                              CROSIER, KATHRYN E.
        9
              (ii) TITLE OF INVENTION: DEVELOPMENTAL TYROSINE KINASES AND
       10
                                       THEIR LIGANDS
             (iii) NUMBER OF SEQUENCES: 16
       12
              (iv) CORRESPONDENCE ADDRESS:
       14
       15
                    (A) ADDRESSEE: NIXON & VANDERHYE P.C.
                                                              ENTERED
       16
                    (B) STREET: 1100 NORTH GLEBE ROAD
      17
                    (C) CITY: ARLINGTON
      18
                    (D) STATE: VIRGINIA
      19
                    (E) COUNTRY: U.S.A.
      20
                   (F) ZIP: 22201-4714
      22
              (V) COMPUTER READABLE FORM:
      23
                   (A) MEDIUM TYPE: Floppy disk
      24
                   (B) COMPUTER: IBM PC compatible
      25
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      26
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
      28
             (vi) CURRENT APPLICATION DATA:
 C--> 29
                   (A) APPLICATION NUMBER: US/09/985,675
 C--> 30
                   (B) FILING DATE: 05-Nov-2001
      36
                   (C) CLASSIFICATION:
     33
            (vii) PRIOR APPLICATION DATA:
                   (A) APPLICATION NUMBER: US 08/505,241
     35
                   (B) FILING DATE: 16-AUG-1995
     38
           (viii) ATTORNEY/AGENT INFORMATION:
     39
                  (A) NAME: MITCHARD, LEONARD C.
     40
                  (B) REGISTRATION NUMBER: 29,009
     41
                  (C) REFERENCE/DOCKET NUMBER: 175-19
            (ix) TELECOMMUNICATION INFORMATION:
     43
     44
                  (A) TELEPHONE: (703) 816-4000
     45
                  (B) TELEFAX: (703) 816-4100
     48 (2) INFORMATION FOR SEQ ID NO: 1:
     49
             (i) SEQUENCE CHARACTERISTICS:
     50
                  (A) LENGTH: 874 AMINO ACIDS
     51
                  (B) TYPE: AMINO ACID
C--> 52
                  (D) TOPOLOGY: LINEAR
     53
            (ii) MOLECULE TYPE: PROTEIN
C--> 54
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
    55 Met
               Gly Trp Pro Gly Leu Arg Pro Leu Leu Leu
    56
                                                             Ala Gly
       1
                              5
    57 Leu
               Ala Ser Leu Leu Pro Gly Ser Ala Ala
                                                     10
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DATE: 02/22/2002 TIME: 09:00:06

					,	- W
58	15			20		
59 Leu	Lys Lei	Met Gl	/ Ala			25
60		30	, ATC			Val Ser
61 Gln	Gly Glr		Tera	3.		
62 40	-1 -1	· IIO vai	-	1 -	s Ser Val	Glu Gly
63 Met	Glu Asp	Dro Acm	45		50	_
64	55 55	4	Ile	His Trp Met	Lys Asp	Gly Thr
65 Val				60		65
66	Val Gln			Gln Val Ser	: Ile Ser	Ile Ser
67 Glu	772 - 0	70			75	201
68	His Ser	Trp Ile	Gly	Leu Leu Ser	Leu Lys	Ser Val
	80			85	-12	90
69 Glu	Arg Ser	Asp Ala	Gly	Leu Tyr Trp	Cys Gln	
70		95		100	-	Val Lys
71 Asp	Gly Glu	Glu Thr	Lys			m
72 105			110	Joz OIII		Trp Leu
73 Thr	Val Glu	Gly Val	Pro	Phe Phe Thr	115	_
74	120	-		125	Val Glu	Pro Lys
75 Asp	Leu Ala	Val Pro	Pro	Asn Ala Pro	_, _,	130
76		135	110	ASII ALA PIO		Leu Ser
77 Cys	Glu Ala	Val Gly	Pro	Dro Cl. D	140	
78	145		FIO	Pro Glu Pro	Val Thr	Ile Tyr
79 Trp	Trp Arg	Gly Leu	mb w	150		155
80		160	Thr	Lys Val Gly	Gly Pro	Ala Pro
81 Ser	Pro Ser			165		
82 170	110 001	Val Leu	Asn	Val Thr Gly	Val Thr	Gln Arg
83 Thr	Glu Phe	C	175		180	_
84	185	Ser Cys	Glu	Ala Arg Asn	Ile Lys	Gly Leu
85 Ala		_		190		195
86	Thr Ser	Arg Pro	Ala	Ile Val Arg	Leu Gln	Ala Pro
87 Pro	31- 31	200			205	,
88	Ala Ala	Pro Phe	Asn	Thr Thr Val	Thr Thr	Ile Ser
89 Ser	210			215		220
90	Tyr Asn	Ala Ser	Val	Ala Trp Val		Ala Asp
	_	225		230	317	ara Asp
91 Gly	Leu Ala	Leu Leu	His	Ser Cys Thr	Val Gln v	Val Ala
92 235			240	•	245	var HIA
93 His	Ala Pro	Gly Glu	Trp	Glu Ala Leu		7-1 17-1
94	250			255	TILG VAI	Val Val
95 Pro	Val Pro	Pro Phe	Thr	Cys Leu Leu	Ara Aan T	260
96		265		1 Lea	Arg Asn I	Leu Ala
97 Pro	Ala Thr	Asn Tyr	Ser	Leu Arg Val		_
98	275	-		280		la Asn
99 Ala	Leu Gly	Pro Ser	Pro	Tyr Gly Asp		85
100	_	290			Trp Val P	ro Phe
101 Gln	Thr Lys	Gly Leu	Ala	295		
102 300	1	<u>7</u> 1300	305	Pro Ala Arg		Gln Asn
103 Phe	His Ala	Ile Arg		3	310	
104	315	AIG	Thr	Asp Ser Gly	Leu Ile	Leu Glu
105 Trp	Glu Glu	Val Ile	D== -	320		325
106	014		Pro	Glu Asp Pro	Gly Glu	Gly Pro
		330			335	_

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				-		–				
107 Leu 108	Gly Pr	o Tyr L	ys Let	ı Se	r Trj	p Val	G1:	n Glu	ı As	n Gly
	340			34	5				35	
109 Thr 110	Gln As	sp Glu L 355	eu Met	: Va	l Gl	u Gly		r Arg		a Asn
111 Leu	Thr As		sp Pro		n T	360				
112 365		rp m	370		и тАз	s Asp	Lei	ı Ile		u Arg
113 Val	Cys Al	a Ser As						375		
114	38		sn Ala	. 116		/ Asp	Gl ⁷	y Pro	Tr	p Ser
115 Gln	Pro Le		.1 0	_	385					390
116	110 10			Se	r His	Asp	His	8 Ala	Gl	y Arg
117 Gln	Gly Pr		95	_			400			
118	405	O Pro Hi	ls Ser			Ser	Trp	> Val	Pro	o Val
119 Val	Leu Gl	u Wal T-		410					415	5
120	Dea Gi		u Thr	Ala	1 Leu	Ile	Thr	Ala	Ala	Ala
121 Leu	Ala Le	420	_			425				
122 430	WIG THE	ı Ile Le		Arg	Lys	Arg	Arg	Lys	Glu	Thr
123 Arg	Pho Cla	- 01 -1	435					440		
124	Phe Gly		a Phe	Asp	Ser	Val	Met	Ala	Arg	Gly
125 Glu	445 Dro 31-				450					455
126	Pro Ala			Arg	Ala	Ala	Arg	Ser	Phe	Asn
127 Arg	Cl., 3	46					465			
128 Alg	Glu Arg	Pro Gl	u Arg		Glu	Ala	Thr	Leu	Asp	Ser
129 Leu		_		475					480	
130	Gly Ile		p Glu	Leu	Lys	Glu	Lys	Leu	Glu	Asp
131 Val	T #1	485				490	_			F
132 495	Leu Ile	Pro Gl	ı Gln	Gln	Phe	Thr	Leu	Gly	Ara	Met
132 495 133 Leu	a 3 -		500					505	5	
134	Gly Lys	_	ı Phe	Gly	Ser	Val	Arg	Glu	Ala	Gln
135 Leu	510				515		_		-,	520
136	Lys Gln	1		Ser	Phe	Val	Lys	Val	Ala	Val
130 137 Lys		525					530			
137 Lys 138	Met Leu	Lys Ala	Asp	Ile	Ile	Ala	Ser	Ser	Asp	Tle
139 Glu	535	_		540					545	
140	Glu Phe	Leu Arg	Glu	Ala	Ala	Cys	Met	Lys	Glu	Phe
140 141 Asp	***	550				555		-		
141 ASP 142 560	His Pro	His Val		Lys	Leu	Val	Gly	Val	Ser	Len
142 360 143 Arg	G	_	565					570		
143 AIG	Ser Arg	Ala Lys	Gly	Arg	Leu	Pro	Ile		Met	Val
144 145 Ile	575				580					585
145 116	Leu Pro	Phe Met	-	His	Gly A	Asp	Leu	His	Ala	
140 147 Leu		590				_	595			2 110
147 Leu 148	Leu Ala	Ser Arg	Ile	Gly	Glu <i>i</i>	Asn	Pro :	Phe	Asn	T.e.u
	600			605					610	пец
149 Pro	Leu Gln	Thr Leu	Val	Arg	Phe N	Met	Val A	Asp	Ile	λ1 >
150		615				520		ı.c.p		nia
151 Cys	Gly Met	Glu Tyr	Leu	Ser :			Asn I	Phe	Ile	ui.
152 625			630			- ,		335	-16	HITP
153 Arg	Asp Leu	Ala Ala		Asn (Cys M	let '	Leu A		C1	N a m
154	640		_		545		u r	4	Glu i	
155 Met	Thr Val	Cys Val	Ala	Asp I		Sly '	Leu S	er	Arg]	550
				-		•			9	-y

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/985,675

DATE: 02/22/2002 TIME: 09:00:06

Input Set : N:\Crf3\RULE60\09985675.raw Output Set: N:\CRF3\02222002\I985675.raw

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	157	T10	Ф			655					660	0		
	158	Tie		s Ser	GΙΣ	/ Asp	Tyr	Туз	Arg	g Gln	Gly	y Cys	Ala	a Ser
	159	Luc	665			_		670				_	675	
	160	цуз	тег	ı Pro		Lys	Trp	Leu	ı Ala	a Leu	Glu	ı Ser		ı Ala
	161 2	Acn	7 ~ ~	т	680					685				
	162		ASI	Leu	Tyr	Thr	Val		Ser	Asp	Val	Trp	Ala	Phe
	163		17-1	ml			695					700		-
	164	3 ± Y	val	Thr 705	мет	Trp	Glu	Ile		Thr	Arg	r Gly	Gln	Thr
	165 E	2ro	Пута		0.1				710					715
	166		TAT	Ala	GIĀ	Ile	Glu	Asn	Ala	Glu	Ile	Tyr	Asn	Tyr
	167 I	en	Tle	Gly	~1	720	_				725	;		-
	168	.cu	730	_	GTÀ	Asn	Arg			Gln	Pro	Pro	Glu	Cys
	169 M	let		Glu	37- 3		_0	735					740	
	170		GIU	GIU		Tyr	Asp	Leu	Met	Tyr	Gln	Cys	Trp	Ser
	171 A	la	λen	Pro	745		_			750				
	172 7		кър	PIO	тĀг	Gln	Arg	Pro	Ser	Phe	Thr	Cys	Leu	Arg
	173 M		Glu	Leu	C1	3	760	_				765		_
	174		GIU	770	GIU	Asn	Ile	Leu		His	Leu	Ser	Val	Leu
	175 s	er		Ser	Cln	7. ~ ~	D	_	775					780
	176	-	- 1111	DCI	GIII	Asp 785	Pro	Leu	Tyr	Ile		Ile	Glu	Arg
	177 A	la	Glu	Gln	Pro		G1	•			790			
	178		795	0111	110	1111	Glu		Gly	Ser	Pro	Glu	Leu	His
	179 C	ys	Gly	Glu	Arg	Sar	So. 77	800	0 1				805	
	180	•	1	Olu	810	261	Ser	ser	Glu		Gly	Asp	Gly	Ser
	181 G	lу	Val	Glv	Ala	Va 1	Gly	C1	71.	815	_			
	182 82			1		val	825	Gly	тте	Pro	Ser		Ser	Arg
	183 ту	/r	Ile	Phe	Ser	Pro	Gly	C1	T	~		830		
	184			835		110	GIY	-		Ser	Glu	Ser	Pro	_
	185 G1	n	Leu		Gln	Gln	Pro	Glu	840	D	_	_		845
	186					850	110	GIU	ser	Pro	Leu	Asn	Glu	Asn
	187 G1	.n	Arg	Leu	Leu		Leu	Gln	Cln	C1	855	_	_	
	188		860					865	GIII	GIY	Leu	Leu	Pro	His
	189 Se		Ser	Cys				005					870	
	192 (2) INF	ORMAT	ION F	OR S	EO IL	NO:	2.						
	193	(i) SEQ	UENCE	CHAI	RACTE	RIST	CS:						
	194		(A) LEN	GTH:	850	AMINO	ACI	os					
	195		(B) TYP	E: Al	ONIN	ACID							
C>			(D) TOP	OLOGY	Z: LI	NEAR							
	197	(ii) MOLI	ECULE	TYPE	: PR	OTEIN	ī						
C>		(xi) SEQU	JENCE	DESC	RIPT	ION:	SEO 1	ED NO): 2.				
	199							~					Ala (71
	200												1	этХ
	201 Le	u	Lys I	Leu 1	Met G	Sly .	Ala	Pro V	al I	.vs	Met 7	Phr	Val S	Sar
	202			5					10	<u> </u>			Yar S	15
	203 Glr	n	Gly G	in i	Pro V	al :	Lys	Leu A		ys :	Ser V	/al	Glu G	
	204	_				20				-	25		Jan G	- T A
	205 Met 206	-	Glu A	sp I	Pro A	sp :	Ile :	His T	rp M	let]	Lys A	ga	Gly T	'hr
2	.00		30					35				•	40	

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/985,675

DATE: 02/22/2002 TIME: 09:00:06

207 **- 1										
207 Val 208	Val Gl	45		Gl	n Va	l Ser		e Ser	Ile	Ser
209 Glu 210 55	His Se	r Trp Ile	e Gly 60		u Le	u Ser	Leu	Lys 65		Val
211 Glu 212	Arg Se:)	a Gly	Le	u Ty 7	r Trp 5	Cys	Gln		Lys 80
213 Asp 214	Gly Gl	85	-	Il	e Se	r Gln	Ser 90	Val	Trp	Leu
215 Thr 216	Val Glu 95	1		Phe 100		e Thr	Val	Glu	Pro 105	Lys
217 Asp 218	Leu Ala	110) Pro	Ası	n Ala	Pro 115	Phe	Gln		Ser
219 Cys 220 120	Glu Ala	-	Pro 125	Pro	o Glu	ı Pro	Val	Thr 130	Ile	Tyr
221 Trp 222	Trp Arg	-		Lys	Va]	l Gly	Gly	Pro	Ala	Pro 145
223 Ser 224	Pro Ser	150		Val	. Thi	Gly	Val 155	Thr	Gln	Arg
225 Thr 226 227 Ala	Glu Phe	-		Ala 165		Asn	Ile	Lys	Gly 170	Leu
228	Thr Ser	175	Ala	Ile	· Val	Arg 180	Leu	Gln	Ala	Pro
229 Pro 230 185 231 Ser	Ala Ala		Asn 190			Val	Thr	Thr 195	Ile	Ser
232 233 Gly	Tyr Asn 200		Val	Ala	Trp 205	Val	Pro	Gly	Ala	Asp 210
234 235 His	Leu Ala	Leu Leu 215	His			Thr	Val 220	Gln	Val	Ala
236 237 Pro	Ala Pro 225 Val Pro	Gly Glu	Trp	230		Leu	Ala	Val	Val 235	Val
238 239 Pro	Ala Thr	Pro Phe 240	Thr			Leu 245	Arg	Asn	Leu	Ala
240 250 241 Ala		Asn Tyr	Ser 255			Val	Arg	Cys 260	Ala	Asn
242 243 Gln	Leu Gly 265 Thr Lys	Pro Ser	Pro		Gly 270	_	Trp		Pro	Phe 275
244 245 Phe	His Ala	Gly Leu 280 Ile Arg	Ala		Ala		Ala 285		Gln /	Asn
246 247 Trp	290 Glu Glu	Val Ile	Thr	295		Gly	Leu		Leu (Glu
248 249 Leu	Gly Pro	305				Pro 310			Gly 1	?ro
250 315 251 Thr	Gln Asp	Tyr Lys Glu Leu	Leu 320	Ser				325	Asn (-
252 253 Leu	330 Thr Asp		Met		335	_	Thr A	_	Ala A	Asn 840
254 255 Val	Cys Ala	Trp Asp		Gln			Leu] 350		Leu A	rg
	Clo WIG	Ser Asn	Ala	Ile	Gly	Asp	Gly E	ro	Trp S	er



VERIFICATION SUMMARY PATENT APPLICATION: US/09/985,675

DATE: 02/22/2002 TIME: 09:00:07

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  L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
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